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(71) Applicant (for all designated States except US): HER MAJESTY IN RIGHT OF CANADA as represented by THE MINISTER OF AGRICULTURE AND AGRI-FOOD CANADA [CA/CA]; Eastern Cereal & Oilseed Research Centre, K.W. Neatby Building, Ottawa, Ontario K1A OC6 (CA).

(72) Inventors; and

(75) Inventors/Applicants (for US only): MIKI, Brian [CA/CA]; 1876 Dorset Drive, Ottawa, Ontario K1H 5V1 (CA). GI-JZEN, Mark [CA/CA]; 848 Princess Avenue, London, Ontario N5W 3M4 (CA). MILLER, Shea [CA/CA]; 69 Kenilworth Street, Ottawa, Ontario K1Y 3Y5 (CA). BOW- MAN, Lu-Ann [CA/CA]; R.R. #3, Listowel, Ontario N4W 3G8 (CA). BATCHELOR, Anthea [GB/CA]; 45 Alexander Street, Ottawa, Ontario K1M 1N1 (CA). HU, Ming [CA/CA]; 279 Kundson Drive, Kanata, Ontario K2K 2N8 (CA). BOUTILIER, Kim [CA/NL]; Groen van Prinsterstraat 87, NL-6702 CP Wageningen (NL).

- (74) Agents: SECHLEY, Konrad, A. et al.; Gowling, Strathy & Henderson, Suite 2600, 160 Elgin Street, Ottawa, Ontario K1P 1C3 (CA).
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(57) Abstract

The present invention is directed to isolated genomic sequences that are differentially expressed within seed-coat tissues. These DNAs are expressed within the inner integument, thick, or thin, walled parenchyma, endothelium, palisade, hourglass, or stellate parenchyma cells of the seed-coat. Furthermore, this invention relates to promoter regions obtained from genomic sequences that are differentially expressed in seed-coat tissues, and their use for directing seed-coat specific expression of genes of interest within transformed plant cells or plants. Exemplified promoters include those obtained from the differential screening of a seed-coat library and cryptic promoters obtained using T-DNA tagging using a promoterless marker gene.

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